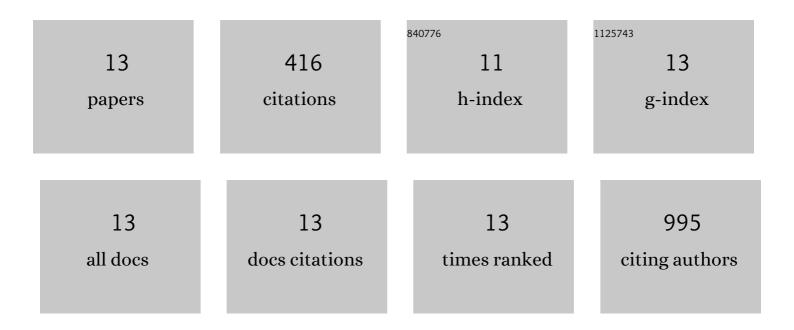


List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5810634/publications.pdf Version: 2024-02-01



#	Article	lF	CITATIONS
1	NF-κB Gene Signature Predicts Prostate Cancer Progression. Cancer Research, 2014, 74, 2763-2772.	0.9	99
2	Loss of FOXA1 Drives Sexually Dimorphic Changes in Urothelial Differentiation and Is an Independent Predictor of Poor Prognosis in Bladder Cancer. American Journal of Pathology, 2015, 185, 1385-1395.	3.8	60
3	Coupled analysis of gene expression and chromosomal location. Genomics, 2005, 85, 401-412.	2.9	47
4	LIM Domain Only-2 (LMO2) Induces T-Cell Leukemia by Two Distinct Pathways. PLoS ONE, 2014, 9, e85883.	2.5	46
5	FOXA1 deletion in luminal epithelium causes prostatic hyperplasia and alteration of differentiated phenotype. Laboratory Investigation, 2014, 94, 726-739.	3.7	39
6	SPARCL1 suppresses metastasis in prostate cancer. Molecular Oncology, 2013, 7, 1019-1030.	4.6	32
7	Strategy for encoding and comparison of gene expression signatures. Genome Biology, 2007, 8, R133.	8.8	24
8	Identification of Genes Required for Enzalutamide Resistance in Castration-Resistant Prostate Cancer Cells <i>In Vitro</i> . Molecular Cancer Therapeutics, 2021, 20, 398-409.	4.1	17
9	Evaluation of public cancer datasets and signatures identifies TP53 mutant signatures with robust prognostic and predictive value. BMC Cancer, 2015, 15, 179.	2.6	15
10	A Murine Model of K-RAS and β-Catenin Induced Renal Tumors Expresses High Levels of E2F1 and Resembles Human Wilms Tumor. Journal of Urology, 2015, 194, 1762-1770.	0.4	15
11	Web-based interrogation of gene expression signatures using EXALT. BMC Bioinformatics, 2009, 10, 420.	2.6	11
12	Identification of a gene-expression predictor for diagnosis and personalized stratification of lupus patients. PLoS ONE, 2018, 13, e0198325.	2.5	7
13	A Data Similarity-Based Strategy for Meta-analysis of Transcriptional Profiles in Cancer. PLoS ONE, 2013, 8, e54979.	2.5	4